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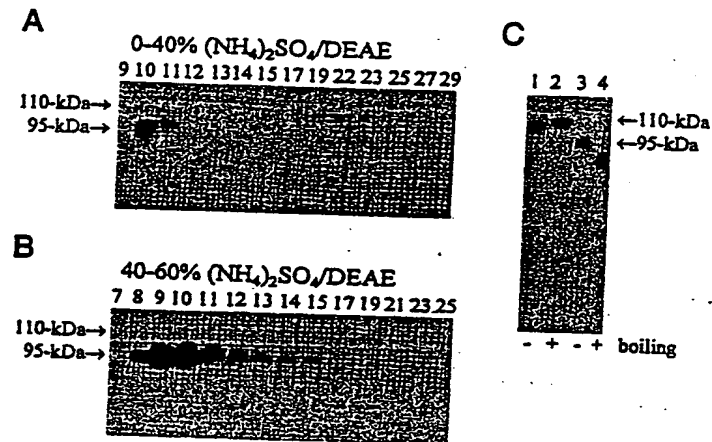


FIG. 1

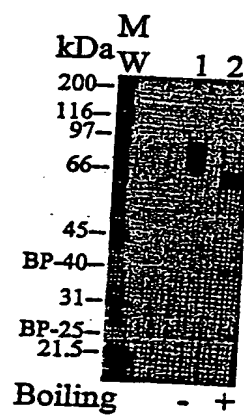


FIG. 2

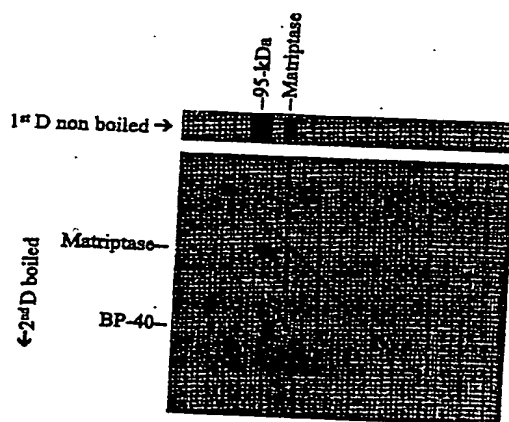


FIG. 3

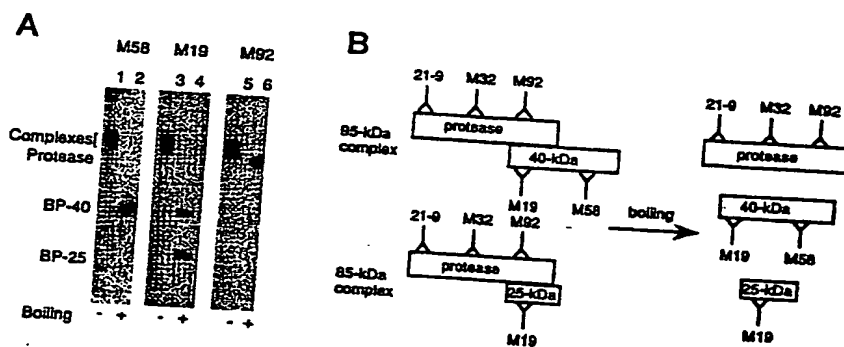


FIG. 4

1 MAPARTMARARLAPAGTPAVALWLLCTLGLOGTOAGPPPA
 41 PPGLPAGADCLNSFTAGVPGFVLDTNASVNGATFLESPT
 81 VRRGWDCVRACCTTQNCMLALVELQPDREGDAIAACFLIN
 121 CLYEQNFVCKFAPREGFINYLTRVYRSYQLRTQGFEGGS
 161 GIPKAWAGIDLKVOPOEPLVLKQVENTDWRLLRGDTDVRV
 201 ERKDPNOVELWGLKEGTYLQTLTVTSSDHPEDTANVTVTV
 241 LSTKQTEDYCLASNKVGRCRGSFPRWYDPTQICKSFVY
 281 GGCLGNKNYLRREECILACRGVOGFSMERHPVCSGTQ
 321 PTQFRCSNGCCIDSFLECDTFCNCPDASDEAAACEKYSGP
 361 DELQRIHFPSDKGHCVDLPDTGLCKESI PRWYNYFFSEHC
 401 ARPTYGGCYGNKNNFEEQQCLESCRGISKKDVFGRLREI
 441 PIPSDGSMEMAVAVFLVICIVVVAILGYCFFKNQRKDFH
 481 GHHHPPTPASSTVSTTEDTEHLVYNHTTRPL

FIG. 5

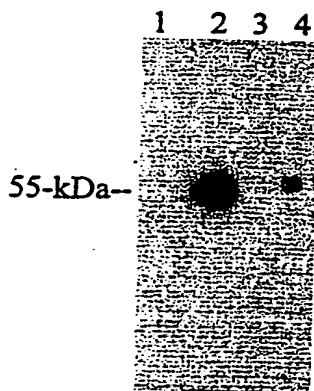


FIG. 6

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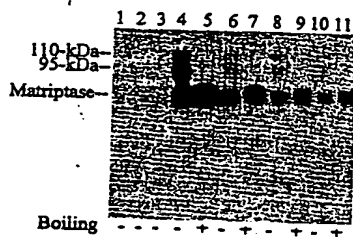


FIG. 7

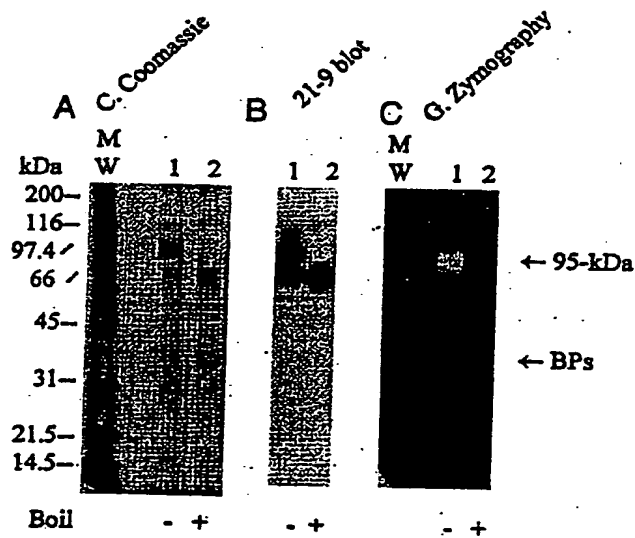


FIG. 8

-357 CGCTGGGGTGTCTGCGACGCGCTGCTGATCGGCTCTCTTGTTCTTGCTGGGATCGGTTCTCGTTCGTCATTTCGATACCGG
 -270 GAGTCGCTGTGCCAAGAGTCTTCAATGGCTACATGAGGATCAAAATGAGAATTTTGTGGATGCCTACGAGAACCTCAACTCCACTGAG
 -180 TTGTAAAGCTGGCCAGAGGTGAAGGACCGCTGAAGCTGTACAGCGGAGTCCCAATCTGGGCCCTACAGCAAGGATCGGCT
 -90 GTGACGGCTTTCAGGACGGGACGCTGATCGCTCAAGCTGTGGTCTGAGTTCAGCATCCCGACACCTGGTGGAGCGGCGCGCGCTC
 1 ATGGCCAGAGCGGTAGTCACTGCGCCGCGGGCGCGCTCCTCAAGTCTTGTGTACCTCAGTGGTGGCTTCTCCACCGAG
 1 M A E E R V V K L P P R A R S L K S F V V T S V Y A F T P D
 91 TCEAAAACAGTACAGGAGACCCAGGACAACAGCTGCAGCTTTGGCTGACGCGCGCGGTGTGAGCTGATGCGCTTACCACGCGCGCGG
 31 S K T V O R T O D N S C S F G L H A R S V E L H R F T T P G
 181 TTCCCTGACAGCCCTACCCGCTCATGCCGCTGCCAGTGGCGCTGCGGGGACCCGAGCTAGTGTGAGCCTACCTTCCGCGAG
 61 F P D S P Y P A H A R C Q W A L R G G D A D S V L S L T F R S
 271 TTGACCTTGGCTCTGCGACGAGCGCGGACGCTGGTACGGTGTACAACACCTGAGCCCCATGGAGCCCAACGCGCTGGTGCAG
 91 F D L A S C T C G D E R G S D L Y T V Y N T L S P H E P H A L V O
 381 TTGTGTGGCACTACCTCCCTCTCAACCTGACCTTCCACTCTCCCAACAGCTCTGCTCATCACACTGAAACCAACACTGAGCGG
 121 L C G T Y P P S Y N L T F H S S Q N V L L I T L I T N T E R
 451 CGGCATCCCGCTTTCAGGCCACCTTCTTCCAGCTGCTAGGATGAGCAGCTGTGGAGGCGGCTTACGTAAGAGCCAGGGGACATTCAAC
 151 R H P G F E A T F F C O L P R H S S C G G R L R K A O G T F N
 541 AGCCCCCTACTACCGAGCCACTACCCCAACATGACTGCACATGGAACACTGAGGTGCCCAACACAGCATGTGAAGTGCAGCTTC
 181 S P Y Y P G H Y P P N I D C T W N I E V P P N N Q H V K V R F
 830 AAATTCTTCTACCTGCTGGAGCCCGCGCTGCTCGGGGACCTGCCCCAAGGACTACGTGGAGATCAATGGGAGAAATACTGCGGAGAG
 211 K F F Y L L E P G V P A G T C P K D Y V E I N G E K Y C G E
 721 AGGTCGAGTTCGTCGTACCCAGCAACAGCAACAGATCACAGTTCGGTTCTCACTCAGTTCAGCTACACCGACCGGCTTCTTAGCT
 241 R S O F V Y T S N S N K I T V R F H S D S Q S Y T D T G F L A
 811 GAATACCTTCTCTACGACTCCAGTGACCCATGCCCGGGGAGTTCAGCTGCGGACGGGCGGTGTATCGGAAGGAGCTGCCCTGTGAT
 271 E Y L S Y D S S O P C G G O F T C R T G R C I R K E L R C D
 901 GGCTGGCGGACGACCCAGCAACAGCATGAGCTCAACTGCAATGCGACGCGGGGACAGTTCAGCTGCAAGAACAGTTCTGCAAG
 301 G W A D C T D H S D E L N C S C D A G H O F T C K N K F C K
 991 CCCCCTCTTTCGGCTTCCGACAGTGTGAACGACTGCGGAGACAACAGCAGCAGCAGCGGTGCACTGTCCGCTCCAGCTTTCAGGTGT
 331 P L F V V C D S V N D C G G D N S D E O G C S C T G A T O T F R C
 1081 TCCAGTGGGAAGTCCCTCTCGAAAGCAGCAGCTGCAATGGGAAGGAGCTGTGGGACCGGTCGACAGGCGCTCTGCCCCAAGGTG
 361 S N G K C L S K S S Q C N G K D D C G D G S D E A S C P K V
 1171 AACGTCGTCACCTTGACCAACACACCTACCGCTGCTCTCAATGGGCTTCTGCTGAGCAAGGGCAACCTGAGTGTGACGGGAAGGAGGAC
 391 N V Y T C T K H T Y R C L H G L C L S K G N P E C D G K G A
 1261 TGTACCGCGCTCAGATGAGAAGGACTGCGACTGTGGGCTGCGGTCACTCAGGAGACAGGCTCGTGTGTTGGGGGACGAGATCGCGAT
 421 C S D G S D E K D C D C G L R S F T R O A R V V G G T D A D
 1351 GAGGCGAGTGGCCTTGGGAGTAAAGCTGATGCTTGGGCGAGGGACATCTGCGGTCTCTCCATCTCTCCCAACTGGCTGGCT
 451 E G E W P M O V S L H A T C G G H I C G A S L I S P N W L Y
 1441 TCTGCGGCACTGCTACATCGATTACAGAGGATTCAGGTACTCAGACCCACGCGAGTGGAGCGGCTTCTGGGCTTGACGACGACAGAG
 481 S A A H C Y I D D R G F R Y S D P T Q W T A L L G L H D Q S
 1531 CAGCGACGCGCCCTGGGCTGAGGAGCGGCTCAAGCGCATCATCTCCACCCCTTCTTCAATGACTTCACCTTCGACTATGATCATC
 511 Q R S A P G V O E R R L K R I S H P P F N D F T F O D I
 1621 CGCCTGCTGGAGCTGAGAAAAACCGCAGAGTACAGCTCCATGGTGGGCGCATCTGCGCTGCGGAGCTCCCATGTCTTCCCTGCCGGC
 541 A L L E L E K P A E Y S S H V R P I C L P D A S H V F P A G
 1711 AAGGCCATCTGGGCTCAGGCGTGGGACACACCCAGTATGAGGAGCATCGCGCGCTGATCTGCAAAAGGGTGAGATCGCGCTCAAC
 571 K A I W V J G W G H T O Y G G S T A L I L O K G E I R V I N
 1801 CAGACCACCTGCGAGAACCTCTGCCGACGAGATCAGCGCGGCGATGATGTGCTGGGCTTCTCAGCGCGCGGCTGAGCTCTGCCAG
 601 Q T T C E N L L P O O I T P R N H C V G G F L S G G G V D S C O
 1891 GGTGATTCGGGGGACCCCTGTCCAGCTGGAGGCGGATGGCGGATCTCCAGGCGGCTGGTGGAGCTGGGAGACCGCTGCGCTCAG
 631 G D S G P L S S V E A D R I F O A G V V S W G D G C A O
 1981 AGGAACAAGCAGCGGTGTACACAAGGCTCCCTCTGTTTGGGAGTGAATCAAGAGAACACTGGGATATAGGGGCGGGGGCACCCAAA
 681 R N K P G V Y T R L P L F R A D V I K E N T G V ...
 2071 TGTGTACACTCGGGGGCCACCATCTGCTACCCCACTGTGCAGCTTCAGGCTGGAGACTGGACCTGACTGCACCGCGCCCCCAAC
 2181 ACATCATCTGTAACCTCAATCTCCAGGCTCAAAATGCTAGAAAACTCTCTCTTCTCAGCTTCAAGTGGAGCTGGGAGGTAG
 2251 AAGGGAGAGACTGGGTTCTACTGACCACTGGGGGCAAGGTTTGAAGACAGAGCTTCCCGGCGACCCCAAGCTGCGGAGG
 2341 CCGGTTGTGATATGCTGCTCCCTTGTCTGAAGGACGCGGAGACGAGCTTGGAGCTTCAAGTGAAGGTTGGGCTGCGG
 2431 ATCTGGGCTTGGGCTTGGGCGACGCTCTGTAGGAGAACCCAGCTCGGAGGAGCTTGGAAACAGCAGGCTTGAAGACTGAAAAATGG
 2521 TTACCAAGCTCCGAGGTGACTTCAGTGTGTATGTGTGAATGAGTAAACATTTTATTTCTTTTAAAAAATGAA

FIG. 9

205060-00000000

Matrilptase Enterokinase THPRS2 Sb-sbd Hepsin Factor XI Plasminogen Trypsin Chymotrypsin	QCL-RSFT	Q--ARVGGT	DADEGECPW	VSLHALG--Q	---GHICGAS	LISPNWLVSA	ARVYIDORG
	SEPK-KLAA	DITPKIVGGS	NAKEGAWPW	VGLY-YG--G	---RLLCGAS	LVSSDWLVSA	ARVYIDORG
	ALGV-NLSS	RQS-RIVGGE	SALPGAWPW	VSLHVQNV--	---HYCGGS	IITPENIVTA	ARVYIDORG
	ECVPTLA--	RQETRIVGCK	SAFGRWPW	VSVRRYSFFG	FSSTHRCGGA	LINENWIATA	ARVYIDORG
	DGR-R--KL	PVD-RIVGGR	DAFGRWPW	VSLRYDTG--	---HLICGGS	LLSGDWLVTA	ARVYIDORG
	ECTT-K----	-IKPRIVGCT	ASVRGCPW	VTLHTTSPTO	---RHLCGGS	IIGNOWILTA	ARVYIDORG
	DCKPOVERK	KCPORVVGCC	VAHPNSWPW	VSLRT--RFG	---MHFCGGT	LISPEWLVTA	ARVYIDORG
	VH-----AAPF	DODDKIVGGY	ICEENSVPYO	VSLN--SGY-	---HFCGGS	LISEOWVSA	ARVYIDORG
	IL-----HPVL	SGLSRIVNGE	DAVGSWPW	VSLQDKTGF-	---HFCGGS	LISEDWVTA	ARVYIDORG
Matrilptase Enterokinase THPRS2 Sb-sbd Hepsin Factor XI Plasminogen Trypsin Chymotrypsin	RYSDPTOWTA	FLGLHDSOR	SAPGVQERRL	KRIISHPFF-	-----NDFTE	DYDIALLELE	KPAEYSSHYR
	ANLEPSKWT	ILGLHKSNL	TSPQTPVRLI	DEIVINPHY-	-----NRRRK	DNDIAMHLE	FKVNYTOYIO
	PLNPPWHTA	FAGILROSEH	F--YGAGYOV	OKVISHPNY-	-----DSKTK	NNDIAMHLE	KPLTFNDLVK
	QIRIRGEYD	FSHYOEOLPY	IERG-----V	AKVYHPKY-	-----LTY	EYDIALVLE-	QPLEFAPHVS
	RNRVLSRWV	FAGAVQAQSP	---HGLOLGV	QAVYHGGYL	PRDPNSEN	SNDIALVHLS	SPLPLEYIO
	---PKILRV	YSGILNOSSEI	KE-DTSFFGV	OEIITHDOY-	-----KHAES	GYDIALKLE	TVNYTDSOR
	---RPSYKY	ILGAHOEVNL	-EPH-----V	OEIEVSRFL	-----EPTRK	---DIALKLS	SPAVITOKVI
	KSRIQVRLGE	H-----NIEV	LEGNEOFINA	AKIIRPKY-	-----NSRTL	DNDIALLKLS	SPAVINSRVS
	RTSDVYVAGE	F-----DQGS	DEENIOVLKI	AKVFKNPKF-	-----SILTV	NNDIALLKLS	SPAVINSRVS
Matrilptase Enterokinase THPRS2 Sb-sbd Hepsin Factor XI Plasminogen Trypsin Chymotrypsin	PICLPDASHV	FPAGKAIWYT	GWGHTOYGGT	G-ALILOKGE	IRVINOTICE	NLLP-Q-Q--	-ITPRHMEVG
	PICLPEENOV	FPGRNCSTIA	GWGTVVYOGT	T-ANILQDAD	VPLLSNERCO	QOHP-EYN--	-ITENHICAG
	PVCLPNPQGM	LOPEOLCWIS	GWGATEEKGK	T-SEVLAAR	VLLTETORCN	SRYVDNL--	-ITPAHICAG
	PICLPETDSL	LI-GHNATVT	GWGRLS-EGG	TLPSVLQEV	VPIVSNDCN	SMFMRAGROE	FIPDIFLCAG
	PVCLPAAQOA	LYDGKICTVT	GWGHTOYGGT	Q-AGVLOEAR	VPLISNDVCN	GADFYGNQ--	-IKPKHFCAG
	PICLPKQDR	NVIYTDCTVT	GWGRLKLRD	I-ONTLOKAK	PLVITNECCO	KRYR-GHK--	-ITHKHICAG
	PICLPSPNV	VADRTECFIT	GWGETO-GTF	G-AGLLKEAO	LPVIEKVCN	RYEFLNGR--	-VQSTELCAG
	ASLPTAPP-	-AAGTESLIS	GWGNTLSSGA	DYPDELOCLD	APVLSQACE	A--SYPGK--	-ITNNHFCAG
	AVCLPSADDD	FPAGILCAIT	GWGNTKYNAN	KTPDKLQAAA	LPLLSNAECK	K--SWGRR--	-ITDVHICAG
Matrilptase Enterokinase THPRS2 Sb-sbd Hepsin Factor XI Plasminogen Trypsin Chymotrypsin	FLSGGVDSQ	GDGSGPLSSV	EADGR---IF	QAGVSWGSG	-CAORNPQVY	TRLPLFRDWI	KENTGV
	YEEGIDDSQ	GDGSGPLHCO	ENN-R---VF	LAGVTSFQYK	-CALPNRQVY	ARVSRFTWI	-OSFLH
	FLGNVDSQ	GDGSGPLV--	---TSNNNIW	LIGDTSWGGG	-CAKAYRPGVY	GNVHFTDWI	YRONKANG
	YETGGDSQ	GDGSGPL---	QAQSDGREF	LAGITSWGIG	-CAEALPGVC	TRISKFTPI	LEHYR
	YREGGIDACK	GDGSGPLVCE	DSISRTPRR	LCGIVSWGIG	-CALAUXPGVY	TKVSDFREWI	FOAIKTHSEA
	HLAGGIDDSQ	GDGSGPLVCF	ENK---KYI	LOGVTSWGL	-CAORNPQVY	INNVVEYDWI	LEKTOAV
	FLEGGIDDSQ	GDGSGPLVSN	GE-----	LOGVTSWGL	-CARPNKPGVY	VRVSRFTWI	EGYHRNN
	--ASGVVDSQ	GDGSGPLVCO	KDGA-----	LVGIVSWGSD	-CAKNRPGVY	TKVNYVDWI	KOTIAANS

FIG. 10

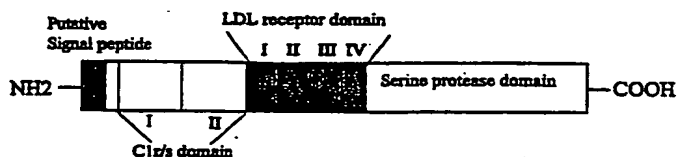
A LDL-receptor type regions

Matriptase (280-314)	PCPG--QFTICRTGRCKIRKELR-CDGWADCTDHSDELNC
(315-351)	SCDAGHQFTICKNKFKCKPLFWV-CDSVNDCGDNDSDEQGC
(352-387)	SCPA-QTERCSNGKCLSKSQ-CCNGKDDCGDGSDEASC
(394-430)	TCTK-HTYRCLNGLCLSKGNPECDGKEDCSGDSDEKDC
Consensus sequences	
LDL-receptor	TC----EF-C--G-CI---W--CD---DC-DGSDE--C
LRP	-C-----F-C---RCIP--W--CDG--DC-D--SDE--C
Perlecan	PC-P-EF-C-----C-----CD---DC-D--SDE--C
GP-330	-C-----F-C-----CI-----CDG--DC-DGSDE--C

B C1r/s type region

Mt (1) 42	CSGLHARGVELMRFTTICFFDSPIAHARQWALRGDADSVLSLTFRS--FDLASCDSGLVT
Mt (2) 168	CGGRLRKAQ-ET--FNSHYFG-HYPPNIDITWIEVPNNQHVKVR-FKFFYLLEPGVPAGT--D
C1r (2) 193	CSSELYTEASQY--ISSLEYPR-SYPPDLRCNYSIRVERGLTLHLKFL-EPFDIDD-HQOVH--D
C1s (2) 175	CSGDVFTALIGE--IASPNYPK-PYPENSRCFYQIRLEKGFQVVTLRREDQVEAADSAGN--D
RaRF (2) 185	CDNLFTORTGV--ITSDFPN-PYPKSSECLYTHELEEGFMVNLQFE-DIFDIED-HREVP--D
CSP (2) 181	CSGDVFTALIGE--IASPNYPK-PYPENSRCFYQIRLEKGFQVVTLRREDQVEAADSAGN--D

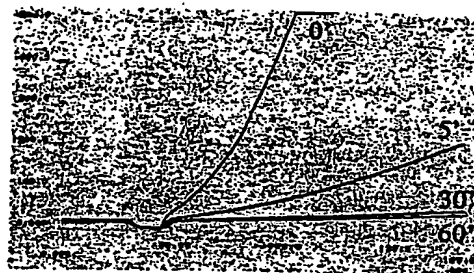
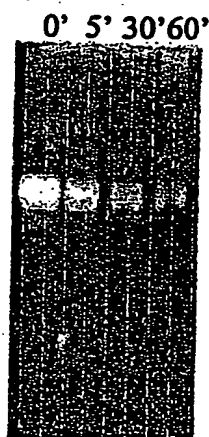
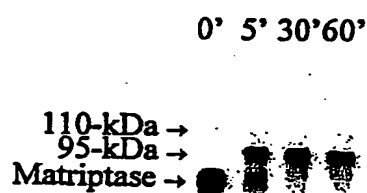
Mt (1) 107	VYNTLS-PHEPHALVQLCQTYFSSYNLTFHSSQVLLITLITNTERRHGF 155
Mt (2) 226	PKDYVEINGE-----YCGER--S-QFVVTNSNKITVRPHSDQSYTDIGF 268
C1r (2) 251	PYDQLQIYANGKNIGFCGXORPP-DLD--TSSNAVOLLFFIDESGDSRGW 298
C1s (2) 235	L-DSLVFVAGDRQFGPYCGHGFFG-PLNIETKSNALDIIFOTDLTGOKKGW 283
RaRF (2) 243	PYDIYIKVGPVKVLGPFCEKAFEPIS--TQSHSVLIUHSNNGENRGW 290
CSP (2) 241	Q-DSLLFAAKNRQFGPFCEGNGFEG-PLTIETHSNITLDIVETDLTEQKKGW 289

FIG. 11**FIG. 12**

A. Immunoblot

B. G. Zymogram

C. Fluorimetric Assay



	Cleavage Rate ¹	Residual Rate
0 min.	725	100%
5 min.	114	18%
30 min.	17	2.3%
60 min.	13	1.8%

FIG. 13

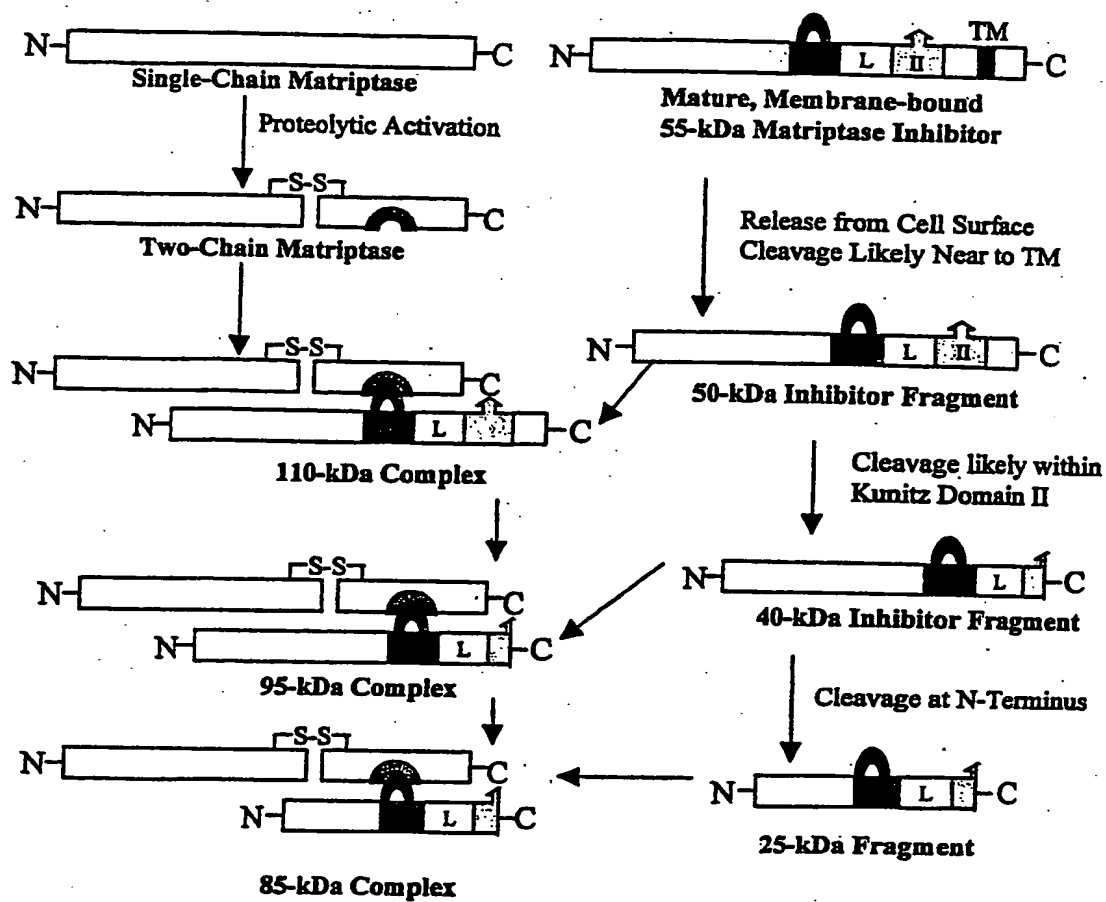


FIG. 14

1 gacgcctgtg agaccgcgca gcggcctcgg ggaccatggg gagcgatcgg gcccgcaagg
61 gcggagggggg cccgaaggac ttcggcgcg gactcaagta caactcccgg cacgagaaaag
121 tgaatggcctt ggaggaaggc gtggagttcc tgccagtcag caactgcaag aaggtggaaa
181 agcatggccc ggggcgctgg gtggtgctgg cagccgtgct gatcgccctc ctcttggtct
241 tgctggggat cggcttcctg gtgtggcatt tgcagtaccg ggacgtgcgt gtccagaagg
301 tcttcaatgg ctacatgagg atcacaatg agaattttgt ggatgcctac gagaactcca
361 actccactga gtttgtaagc ctggccagca aggtgaagga cgcgctgaag ctgctgtaca
421 gcggagtcctt attcctgggc cctaccaca aggagtcggc tgtgacggcc ttcagcgagg
481 gcagcgtcat cgcctactac tgggtctgagt tcagcatccc gcagcacctg gtggaggagg
541 ccgagcgcgt catggccgag gagcgcgtag tcatgctgcc cccgcgggcg cgtccctga
601 agtcctttgt ggtcacctca gtggtggctt tccccacgga ctccaaaaca gtacagagga
661 cccaggacaa cagctgcagc tttggcctgc acgcccgcgg tgtggagctg atgctgttca
721 ccacgcccgg cttccctgac agcccctacc ccgctcatgc ccgctgccag tgggcccctgc
781 ggggggacgc cgactcagtg ctgagcctca ccttcgcag ctttgacctt gcgtcctgcg
841 acgagcgcgg cagcgacctg gtgacggtgt acaacaccct gagccccatg gagccccacg
901 ccctggtgca gttgtgtggc acctaccctc cctcctacaa cctgaccttc cactcctccc
961 agaacgtcct gctcatcaca ctgataacca acactgagcg gcggcatccc ggctttgagg
1021 ccaccttctt ccagctgcct aggatgagca gctgtggagg ccgcttacgt aaagcccagg
1081 ggacattcaa cagcccctac taccaggcc actaccacc caacattgac tgcacatgga
1141 acattgaggt gcccaacaac cagcatgtga aggtgcgctt caaattcttc tacctgctgg
1201 agccggcggt gcctgcgggc acctgccccca aggactacgt ggagatcaat ggggagaaat
1261 actgcgagga gaggtcccag ttcgtcgtca ccagcaacag caacaagatc acagttcgct
1321 tccactcaga tcagtcctac accgacaccg gcttcttagc tgaatacctc tcctacgact
1381 ccagtgaacc atgcccgggg cagttcacgt gccgcacggg gcggtgtatc cggaaggagc
1441 tgcgctgtga tggtggggc gactgcaccg accacagcga tgagctcaac tgcagttgcg
1501 acgcccggca ccagttcacg tgcaagaaca agttctgcaa gcccctcttc tgggtctgcg
1561 acagtgtgaa cgactgcgga gacaacagcg acgagcaggg gtgcagttgt ccggcccaga
1621 ccttcaggtg ttccaatggg aagtgcctct cgaaaagcca gcagtgaat gggaaggagc
1681 actgtgggga cgggtccgac gaggcctcct gcccgaaggt gaacgtcgtc acttgtacca
1741 aacacaccta ccgctgcctc aatgggctct gcttgagcaa gggcaaccct gagtgtgacg
1801 ggaaggagga ctgtagcgac ggctcagatg agaaggactg cgactgtggg ctgcggtcat
1861 tcacgagaca ggctcgtgtt gttgggggca cgatgcgga tgagggcgag tggccctggc
1921 aggtaagcct gcatgctctg ggccaggggc acatctgcgg tgcttccctc atcttccca
1981 actggctggt ctctgcgca cactgctaca tcgatgacag aggattcagg tactcagacc
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2101 tgcaggagcg caggctcaag cgcctcatct cccaccctt cttcaatgac ttcacctcg
2161 actatgacat cgcgctgctg gagctggaga aaccggcaga gtacagctcc atggtgcggc
2221 ccatctgcct gccggacgcc tccatgtct tccctgccgg caaggccatc tgggtcacgg
2281 gctggggaca caccagtat ggaggcactg gcgcgctgat cctgcaaaag ggtgagatcc
2341 gcgtcatcaa ccagaccacc tgcgagaacc tcctgcgca gcagatcag ccgcgcatga
2401 tgtgcgtggg cttcctcagc ggcggcggtg actcctgcca ggggtattcc gggggacccc
2461 tgtccagcgt ggaggcggat gggcggtatc tccaggccgg tgtggtgagc tggggagacg
2521 gctgcgctca gaggaacaag ccaggcgtgt acacaaggct ccctctgttt cgggactgga
2581 tcaaagagaa cactggggta taggggcccgg ggccacccaa atgtgtacac ctgcggggccc
2641 acccatcgtc caccacagtg tgcacgcctg caggctggag actggaccgc tgactgcacc
2701 agcgcgccca gaacatacac tgtgaactca atctccaggg ctccaaatct gcctagaaaa
2761 cctctcgctt cctcagcctc caaagtggag ctgggaggta gaaggggagg acactggtgg
2821 ttctactgac ccaactgggg gcaaagggtt gaagacacag cctccccgcg cagccccaag
2881 ctggggccgag gcgcgtttgt gtatatctgc ctcctctgtc tgtaaggagc agcgggaacg
2941 gagcttcgga gcctcctcag tgaagggtgt ggggctgccg gatctgggct gtggggccct
3001 tggggccacg tcttgaggaa gcccaggctc ggaggaccct ggaaaacaga cgggtctgag
3061 actgaaaatg gtttaccagc tcccagggtc cttcagtggt tgtattgtgt aaatgagtaa
3121 aacattttat ttctttttaa aaaaaaaaa

FIG. 15

1 mgsdrarkgg ggpkdfigagl kynsrhekvngleegveflp vnnvkkvekh gpgrwvvlaa
61 vliglllvll gigflvwhlg yrdvrvqkvf ngymritnen fvdagensns tefvslaskv
121 kdalkllysg vpflgpyhke savtafsegs viayywsefs ipqhlveeae rvmaeervvm
181 lpprarslks fvvtsvvaftp tdsktvqrtq dnscsfglha rgvelmrftt pgfpdspypa
241 harcqwalg dadsvlsltf rsfdlascde rgsdlvtvyn tlpmphepal vqlcgtypps
301 ynltfhssqn vllitlitnt errhpgfeat ffqlprmsc ggrrlkaagt fnspyyppghy
361 ppnidctwni evpnnqhvkv rfkffyllep gvpagtcpkd yveingekyc gersqfvvts
421 nsnkitvrhf sdqsytdtgf laeysydss dpcpgqftcr tgrcirkelr cdgwadctdh
481 sdelncscda ghqftcknkf ckplfwvcds vndcgdnsde qgcscpaqtf rcsngkclsk
541 sqqcngkddc gdgsdeascp kvnvtctkh tyrclnglcl skgnpecdgk edcsdgsdek
601 dcdcglsrft rgarvvggtd adegewpwqv slhalggghi cgaslispnw lvsaahcyid
661 drgfrysdpt qwtaflglhd qsqrsgpgvq errlkriish pffndftfdy diallelekp
721 aeyssmvrpi clpdashvfp agkaiwvtgw ghtqygggtga lilqkgeirv inqttcenll
781 pqqitprmmc vgflsggvds cggdsggpls sveadgrifq agvvswgdgc aqrnkpgvyt
841 rlpfrdwik entgv

FIG. 16

2050E0-EE9E660

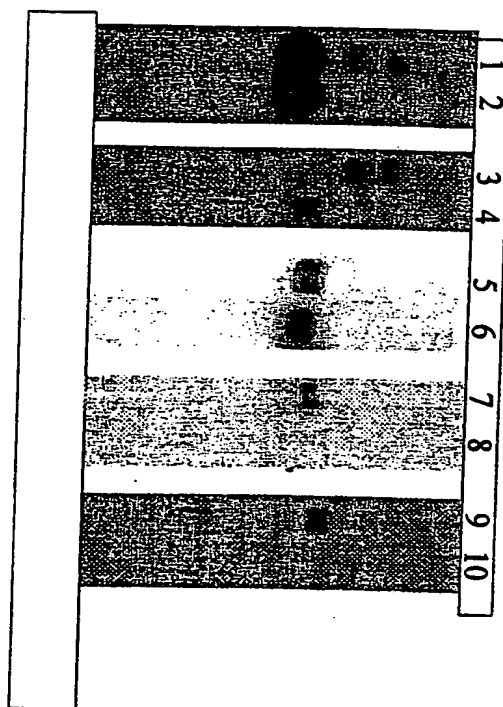


FIG. 17

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